

#5
LMS
3-15-01

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/673,198
 DATE: 01/04/2001
 TIME: 11:31:43

Input Set : A:\Pto.amc
 Output Set: N:\CRF3\01042001\I673198.raw

2 <110> APPLICANT: MIYAKE, Koichiro; HASHIMOTO, Shinichi; MOTOYAMA Hiroaki;
 3 OZAKI, Akio; SEFO, Haruo; KUZAYAMA, Tomohisa; TAKAHASHI, Shunji
 5 <120> TITLE OF INVENTION: A process for producing isoprenoid compounds by
 6 microorganisms and a method for screening compounds with
 7 antibiotic or weeding activity
 W--> 9 <130> FILE REFERENCE:
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/673,198
 C--> 12 <141> CURRENT FILING DATE: 2000-10-12
 14 <150> PRIOR APPLICATION NUMBER: JP98/103101
 15 <151> PRIOR FILING DATE: 1998-04-14
 17 <150> PRIOR APPLICATION NUMBER: JP98/221910
 18 <151> PRIOR FILING DATE: 1998-08-05
 20 <150> PRIOR APPLICATION NUMBER: JP99/035739
 21 <151> PRIOR FILING DATE: 1999-02-15
 23 <160> NUMBER OF SEQ ID NOS: 34
 25 <170> SOFTWARE: PatentIn Ver. 2.0
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 620
 29 <212> TYPE: PRT
 30 <213> ORGANISM: Escherichia coli
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 36 Thr Gln Glu Leu Arg Leu Leu Pro Lys Glu Ser Leu Pro Lys Leu Cys
 37 20 25 30
 39 Asp Glu Leu Arg Arg Tyr Leu Leu Asp Ser Val Ser Arg Ser Ser Gly
 40 35 40 45
 42 His Phe Ala Ser Gly Leu Gly Thr Val Glu Leu Thr Val Ala Leu His
 43 50 55 60
 45 Tyr Val Tyr Asn Thr Pro Phe Asp Gln Leu Ile Trp Asp Val Gly His
 46 65 70 75 80
 48 Gln Ala Tyr Pro His Lys Ile Leu Thr Gly Arg Arg Asp Lys Ile Gly
 49 85 90 95
 51 Thr Ile Arg Gln Lys Gly Gly Leu His Pro Phe Pro Trp Arg Gly Glu
 52 100 105 110
 54 Ser Glu Tyr Asp Val Leu Ser Val Gly His Ser Ser Thr Ser Ile Ser
 55 115 120 125
 57 Ala Gly Ile Gly Ile Ala Val Ala Ala Glu Lys Glu Gly Lys Asn Arg
 58 130 135 140
 60 Arg Thr Val Cys Val Ile Gly Asp Gly Ala Ile Thr Ala Gly Met Ala
 61 145 150 155 160
 63 Phe Glu Ala Met Asn His Ala Gly Asp Ile Arg Pro Asp Met Leu Val
 64 165 170 175
 66 Ile Leu Asn Asp Asn Glu Met Ser Ile Ser Glu Asn Val Gly Ala Leu
 67 180 185 190
 69 Asn Asn His Leu Ala Gln Leu Leu Ser Gly Lys Leu Tyr Ser Ser Leu
 70 195 200 205

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72 Arg Glu Gly Gly Lys Lys Val Phe Ser Gly Val Pro Pro Ile Lys Glu
73      210      215      220
75 Leu Leu Lys Arg Thr Glu Glu His Ile Lys Gly Met Val Val Pro Gly
76 225      230      235      240
78 Thr Leu Phe Glu Glu Leu Gly Phe Asn Tyr Ile Gly Pro Val Asp Gly
79      245      250      255
81 His Asp Val Leu Gly Leu Ile Thr Thr Leu Lys Asn Met Arg Asp Leu
82      260      265      270
84 Lys Gly Pro Gln Phe Leu His Ile Met Thr Lys Lys Gly Arg Gly Tyr
85      275      280      285
87 Glu Pro Ala Glu Lys Asp Pro Ile Thr Phe His Ala Val Pro Lys Phe
88      290      295      300
90 Asp Pro Ser Ser Gly Cys Leu Pro Lys Ser Ser Gly Gly Leu Pro Ser
91 305      310      315      320
93 Tyr Ser Lys Ile Phe Gly Asp Trp Leu Cys Glu Thr Ala Ala Lys Asp
94      325      330      335
96 Asn Lys Leu Met Ala Ile Thr Pro Ala Met Arg Glu Gly Ser Gly Met
97      340      345      350
99 Val Glu Phe Ser Arg Lys Phe Pro Asp Arg Tyr Phe Asp Val Ala Ile
100      355      360      365
102 Ala Glu Gln His Ala Val Thr Phe Ala Ala Gly Leu Ala Ile Gly Gly
103      370      375      380
105 Tyr Lys Pro Ile Val Ala Ile Tyr Ser Thr Phe Leu Gln Arg Ala Tyr
106 385      390      395      400
108 Asp Gln Val Leu His Asp Val Ala Ile Gln Lys Leu Pro Val Leu Phe
109      405      410      415
111 Ala Ile Asp Arg Ala Gly Ile Val Gly Ala Asp Gly Gln Thr His Gln
112      420      425      430
114 Gly Ala Phe Asp Leu Ser Tyr Leu Arg Cys Ile Pro Glu Met Val Ile
115      435      440      445
117 Met Thr Pro Ser Asp Glu Asn Glu Cys Arg Gln Met Leu Tyr Thr Gly
118      450      455      460
120 Tyr His Tyr Asn Asp Gly Pro Ser Ala Val Arg Tyr Pro Arg Gly Asn
121 465      470      475      480
123 Ala Val Gly Val Glu Leu Thr Pro Leu Glu Lys Leu Pro Ile Gly Lys
124      485      490      495
126 Gly Ile Val Lys Arg Arg Gly Glu Lys Leu Ala Ile Leu Asn Phe Gly
127      500      505      510
129 Thr Leu Met Pro Glu Ala Ala Lys Val Ala Glu Ser Leu Asn Ala Thr
130      515      520      525
132 Leu Val Asp Met Arg Phe Val Lys Pro Leu Asp Glu Ala Leu Ile Leu
133      530      535      540
135 Glu Met Ala Ala Ser His Glu Ala Leu Val Thr Val Glu Glu Asn Ala
136 545      550      555      560
138 Ile Met Gly Gly Ala Gly Ser Gly Val Asn Glu Val Leu Met Ala His
139      565      570      575
141 Arg Lys Pro Val Pro Val Leu Asn Ile Gly Leu Pro Asp Phe Phe Ile
142      580      585      590
144 Pro Gln Gly Thr Gln Glu Glu Met Arg Ala Glu Leu Gly Leu Asp Ala

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153 <212> TYPE: PRT
154 <213> ORGANISM: Escherichia coli
156 <400> SEQUENCE: 2
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160 Ala Leu Ser Arg Phe Ile Ala Pro Leu Pro Phe Gln Asn Thr Pro Val
161 20 25 30
163 Val Glu Thr Met Gln Tyr Gly Ala Leu Leu Gly Gly Lys Arg Leu Arg
164 35 40 45
166 Pro Phe Leu Val Tyr Ala Thr Gly His Met Phe Gly Val Ser Thr Asn
167 50 55 60
169 Thr Leu Asp Ala Pro Ala Ala Val Glu Cys Ile His Ala Tyr Ser
170 65 70 75 80
172 Leu Ile His Asp Asp Leu Pro Ala Met Asp Asp Asp Leu Arg Arg
173 85 90 95
175 Gly Leu Pro Thr Cys His Val Lys Phe Gly Glu Ala Asn Ala Ile Leu
176 100 105 110
178 Ala Gly Asp Ala Leu Gln Thr Leu Ala Phe Ser Ile Leu Ser Asp Ala
179 115 120 125
181 Asp Met Pro Glu Val Ser Asp Arg Asp Arg Ile Ser Met Ile Ser Glu
182 130 135 140
184 Leu Ala Ser Ala Ser Gly Ile Ala Gly Met Cys Gly Gly Gln Ala Leu
185 145 150 155 160
187 Asp Leu Asp Ala Glu Gly Lys His Val Pro Leu Asp Ala Leu Glu Arg
188 165 170 175
190 Ile His Arg His Lys Thr Gly Ala Leu Ile Arg Ala Ala Val Arg Leu
191 180 185 190
193 Gly Ala Leu Ser Ala Gly Asp Lys Gly Arg Arg Ala Leu Pro Val Leu
194 195 200 205
196 Asp Lys Tyr Ala Glu Ser Ile Gly Leu Ala Phe Gln Val Gln Asp Asp
197 210 215 220
199 Ile Leu Asp Val Val Gly Asp Thr Ala Thr Leu Gly Lys Arg Gln Gly
200 225 230 235 240
202 Ala Asp Gln Gln Leu Gly Lys Ser Thr Tyr Pro Ala Leu Leu Gly Leu
203 245 250 255
205 Glu Gln Ala Arg Lys Lys Ala Arg Asp Leu Ile Asp Asp Ala Arg Gln
206 260 265 270
208 Ser Leu Lys Gln Leu Ala Glu Gln Ser Leu Asp Thr Ser Ala Leu Glu
209 275 280 285
211 Ala Leu Ala Asp Tyr Ile Ile Gln Arg Asn Lys
212 290 295
215 <210> SEQ ID NO: 3
216 <211> LENGTH: 80
217 <212> TYPE: PRT

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224 Glu Leu Glu Gln Ile Val Thr Arg Leu Glu Ser Gly Asp Leu Pro Leu
225 20 25 30
227 Glu Glu Ala Leu Asn Glu Phe Glu Arg Gly Val Gln Leu Ala Arg Gln
228 35 40 45
230 Gly Gln Ala Lys Leu Gln Gln Ala Glu Gln Arg Val Gln Ile Leu Leu
231 50 55 60
233 Ser Asp Asn Glu Asp Ala Ser Leu Thr Pro Phe Thr Pro Asp Asn Glu
234 65 70 75 80
237 <210> SEQ ID NO: 4
238 <211> LENGTH: 348
239 <212> TYPE: PRT
240 <213> ORGANISM: Escherichia coli
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244 1 5 10 15
246 Asp Glu Tyr Ser Arg Ser Gly Ser Met Gln Tyr Asn Pro Leu Gly Lys
247 20 25 30
249 Thr Asp Leu Arg Val Ser Arg Leu Cys Leu Gly Cys Met Thr Phe Gly
250 35 40 45
252 Glu Pro Asp Arg Gly Asn His Ala Trp Thr Leu Pro Glu Glu Ser Ser
253 50 55 60
255 Arg Pro Ile Ile Lys Arg Ala Leu Glu Gly Ile Asn Phe Phe Asp
256 65 70 75 80
258 Thr Ala Asn Ser Tyr Ser Asp Gly Ser Ser Glu Glu Ile Val Gly Arg
259 85 90 95
261 Ala Leu Arg Asp Phe Ala Arg Arg Glu Asp Val Val Val Ala Thr Lys
262 100 105 110
264 Val Phe His Arg Val Gly Asp Leu Pro Glu Gly Leu Ser Arg Ala Gln
265 115 120 125
267 Ile Leu Arg Ser Ile Asp Asp Ser Leu Arg Arg Leu Gly Met Asp Tyr
268 130 135 140
270 Val Asp Ile Leu Gln Ile His Arg Trp Asp Tyr Asn Thr Pro Ile Glu
271 145 150 155 160
273 Glu Thr Leu Glu Ala Leu Asn Asp Val Val Lys Ala Gly Lys Ala Arg
274 165 170 175
276 Tyr Ile Gly Ala Ser Ser Met His Ala Ser Gln Phe Ala Gln Ala Leu
277 180 185 190
279 Glu Leu Gln Lys Gln His Gly Trp Ala Gln Phe Val Ser Met Gln Asp
280 195 200 205
282 His Tyr Asn Leu Ile Tyr Arg Glu Glu Glu Arg Glu Met Leu Pro Leu
283 210 215 220
285 Cys Tyr Gln Glu Gly Val Ala Val Ile Pro Trp Ser Pro Leu Ala Arg
286 225 230 235 240
288 Gly Arg Leu Thr Arg Pro Trp Gly Glu Thr Thr Ala Arg Leu Val Ser
289 245 250 255

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Input Set : A:\Pto.amc

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291 Asp Glu Val Gly Lys Asn Leu Tyr Lys Glu Ser Asp Glu Asn Asp Ala
292      260      265      270
294 Gln Ile Ala Glu Arg Leu Thr Gly Val Ser Glu Glu Leu Gly Ala Thr
295      275      280      285
297 Arg Ala Gln Val Ala Leu Ala Trp Leu Leu Ser Lys Pro Gly Ile Ala
298      290      295      300
300 Ala Pro Ile Ile Gly Thr Ser Arg Glu Glu Glu Leu Asp Glu Leu Leu
301 305      310      315      320
303 Asn Ala Val Asp Ile Thr Leu Lys Pro Glu Gln Ile Ala Glu Leu Glu
304      325      330      335
306 Thr Pro Tyr Lys Pro His Pro Val Val Gly Phe Lys
307      340      345
310 <210> SEQ ID NO: 5
311 <211> LENGTH: 398
312 <212> TYPE: PRT
313 <213> ORGANISM: Escherichia coli.
315 <400> SEQUENCE: 5
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319 Thr Leu Asp Val Val Arg His Asn Pro Glu His Phe Arg Val Val Ala
320      20      25      30
322 Leu Val Ala Gly Lys Asn Val Thr Arg Met Val Glu Gln Cys Leu Glu
323      35      40      45
325 Phe Ser Pro Arg Tyr Ala Val Met Asp Asp Glu Ala Ser Ala Lys Leu
326      50      55      60
328 Leu Lys Thr Met Leu Gln Gln Gly Ser Arg Thr Glu Val Leu Ser
329      65      70      75      80
331 Gly Gln Gln Ala Ala Cys Asp Met Ala Ala Leu Glu Asp Val Asp Gln
332      85      90      95
334 Val Met Ala Ala Ile Val Gly Ala Ala Gly Leu Leu Pro Thr Leu Ala
335      100      105      110
337 Ala Ile Arg Ala Gly Lys Thr Ile Leu Leu Ala Asn Lys Glu Ser Leu
338      115      120      125
340 Val Thr Cys Gly Arg Leu Phe Met Asp Ala Val Lys Gln Ser Lys Ala
341      130      135      140
343 Gln Leu Leu Pro Val Asp Ser Glu His Asn Ala Ile Phe Gln Ser Leu
344 145      150      155      160
346 Pro Gln Pro Ile Gln His Asn Leu Gly Tyr Ala Asp Leu Glu Gln Asn
347      165      170      175
349 Gly Val Val Ser Ile Leu Leu Thr Gly Ser Gly Gly Pro Phe Arg Glu
350      180      185      190
352 Thr Pro Leu Arg Asp Leu Ala Thr Met Thr Pro Asp Gln Ala Cys Arg
353      195      200      205
355 His Pro Asn Trp Ser Met Gly Arg Lys Ile Ser Val Asp Ser Ala Thr
356      210      215      220
358 Met Met Asn Lys Gly Leu Glu Tyr Ile Glu Ala Arg Trp Leu Phe Asn
359 225      230      235      240
361 Ala Ser Ala Ser Gln Met Glu Val Leu Ile His Pro Gln Ser Val Ile
362      245      250      255

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/673,198

DATE: 01/04/2001

TIME: 11:31:44

Input Set : A:\Pto.amc

Output Set: N:\CRF3\01042001\I673198.raw

L:9 M:201 W: Mandatory field data missing, FILE REFERENCE
L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1054 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 11
L:1055 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1060 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1065 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1070 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1075 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1080 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1085 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1090 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1095 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1100 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1105 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1110 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
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L:1135 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
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VERIFICATION SUMMARY
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Input Set : A:\Pto.amc

Output Set: N:\CRF3\01042001\I673198.raw

L:1275 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1280 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
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L:1290 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1295 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11
L:1300 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:11

Serial Number: 673,198

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☒ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically:
-
- ☒ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
-
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
-
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
-
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included:
-
- ☐ Deleted extra, invalid, headings used by an applicant, specifically:
-
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as
-
- ☐ Inserted mandatory headings, specifically:
-
- ☐ Corrected an obvious error in the response, specifically:
-
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically:
-
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected:
-
- ☐ Other:
-

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.